SEQUENCE LISTING

#5

<110> Crucell Holland B.V.

Houtzager, Erwin

Logtenberg, Ton

de Kruif, Cornelis A.

<120> Chimaeric Phages

<130> 2183-4957US

<140> US 09/882,621

<141> 2001-06-15

<160> 8

<170> PatentIn version 3.1

<210> 1

<211> 30

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer D3 BamHI Forward

<400> 1
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<210> 2

<211> 30

<212> DNA

<213> Artificial sequence

<220>

<223> Primer D3 BamHI Backward

<400> 2
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<210> 3

<211> 33

<212> DNA

<213> Artificial sequence

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<223> Primer g3 minus HindIII Forward

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<210> 4

<211> 31

<212> DNA

<213> Artificial sequence

<220>

<223> Primer g3 minus HindIII Backward

<400> 4
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<210> 5

<211> 31

<212> DNA

<213> Artificial sequence

<220>

<223> Primer g3 ORF NcoI Forward

<400> 5

2

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<210> 6

<211> 31

<212> DNA

<213> Artificial sequence

<220>

<223> Primer g3 ORF XbaI Backward

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<210> 7

<211> 266

<212> DNA

<213> Artificial sequence

<220>

<223> Sequence of the part of the helper phage that surrounds the g3 de
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cttttttttt ggagattttc aacaagcttc tgcgtaataa ggagtcttaa tcatgccagt     180

tcttttgggt attccgttat tattgcgttt cctcggtttc cttctggtaa ctttgttcgg     240

ctatctgcta acttttctta aaaagg                                          266
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<210> 8

<211> 832

<212> DNA

<213> Artificial sequence

<220>

<223> Representation of the helper phage genome deleted for part of the

g3 gene

<400> 8
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gataaaccga tacaattaaa ggctcctttt ggagcctttt tttttggaga ttttcaacgt     180

gaaaaaatta ttattcgcaa ttcctttagt tgttcctttc tattctcact ccgctaagct     240

ttctggttcc ggtgattttg attatgaaaa tatggcaaac gctaataagg gggctatgac     300

cgaaaatgcc gatgaaaacg cgctacagtc tgacgctaaa ggcaaacttg attctgtcgc     360

tactgattac ggtgctgcta tcgacggttt cattggtgac gtttccggcc ttgctaatgg     420

taatggtgct actggtgatt ttgctggctc taattcccaa atggctcaag tcggtgacgg     480

tgataattca cctttaatga ataatttccg tcaatattta ccttcccttc ctcaatcggt     540

tgaatgtcgc ccttttgtct ttcgcgctgg taaaccatat gaattttcta ttgattgtga     600

caaaataaac ttattccgtg gtgtctttgc gtttctttta tatgttgcca cctttatgta     660

tgtattttcg acgtttgcta acatactgcg taataaggag tcttaatcat gccagttctt     720

ttgggtattc cgttattatt gcgtttcctc ggtttccttc tggtaacttt gttcggctat     780

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